PATENT

Docket No.: 28594/38247

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

| Applicants: Shannon et al. | "EXPRESS MAIL" mailing label No.EU 424 299 483 US |
|------------------------------------|--|
| Serial No: 10/049,953 |) Date of Deposit: 17 June 2002 |
| Filed: February 19, 2002 | I hereby certify that this paper (or fee) is |
| Title: Recombinant Subunit Vaccine | being deposited with the United StatesPostal Service "EXPRESS MAIL POST |
| Group Art Unit: To be assigned | OFFICE TO ADDRESSEE" service under 37 C.F.R. §1.10 on the date indicated above |
| Examiner: To be assigned | and is addressed to the Commissioner for Patents Box PCT. Washington, DC 20231 Richard Zimmermann |

STATEMENT PURSUANT TO 37 C.F.R. §§1.825(a) and (b)

Commissioner for Patents Box PCT Washington, DC 20231

Sir:

I hereby state that the contents of the substitute paper copy (pages 1-3 submitted herewith) and the substitute computer readable form of the sequence listing submitted herewith in accordance with 37 C.F.R. §§1.825(a) and (b), are the same and do not introduce new matter into the disclosure of the application.

Respectfully submitted,

06/20/2002 HKAYPAGH 00000029 10049953 01 FC:154

130.00 OP

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By:

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Registration No. 40,994

Attorney for Applicants-

June 17, 2002



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

| Application Serial Number: | 10/049,953 | |
|----------------------------|------------|--|
| Source: | PCTIO | |
| Date Processed by STIC: | 3/5/02 | |

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS. PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual - ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
- 3. Hand Carry directly to: U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
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Revised 01/29/2002

| | 10/01/9 953 |
|-------------------------------------|---|
| ERROR DETECTED | SUGGESTED CORRECTION SERIAL NUMBER: 10/049,953 |
| ATTN: NEW RULES CASES: | PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE |
| 1Wrapped Nucleics Wrapped Aminos | The number/text at the end of each line "wrapped" down to the next line. This may be used in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping." |
| 2Invalid Line Length | The rules require that a line not exceed 72 characters in length. This includes white spaces. |
| 3Misaligned Amino Numbering | The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead. |
| 4Non-ASCII | The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text. |
| 5Variable Length | Sequence(s)contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each each please present having variable length and indicate in the <220>-<223> section that some may be missing. |
| 6PatentIn 2.0 "bug" | A "bug" in Patentln version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, Patentln would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for |
| 7Skipped Sequences (OLD RULES) | Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This requence is intentionally skipped |
| | Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to Include the skipped sequences. |
| 8Skipped Sequences (NEW RULES) | <210> sequence id number <400> sequence id number 000 |
| 9 Use of n's or Xaa's (NEW RULES) | Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents. |
| 10 Invalid <213> Response | Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence |
| 11 Use of <220> | Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or use of capacity and account of genetic material in <220> to <223> section. |
| 12Patentln 2.0 "bug" | Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk. |
| 13Misuse of n | n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide. |
| | AMC/MH - Biotechnology Systems Branch - 08/21/2001 |



Does Not Comply Corrected Diskette Needed

PCT10

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/049,953

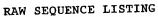
DATE: 03/05/2002 TIME: 14:11:51

Input Set : A:\38247.txt

Output Set: N:\CRF3\03052002\J049953.raw

The type of errors shown exist throughout
the Sequence Listing. Please check subsequent
sequences for similar errors.

| | sequences for sittilial errors. |
|---|---------------------------------|
| 6 <110> APPLICANT: SHANNON ET AL. | |
| 0 /120 TITLE OF INVENTION: RECOMBINANT SOUGHT (MODELL | |
| | |
| - 15 (140) CUPPENT APPLICATION NUMBER. OD/ 20/ | |
| | |
| C> 15 <141> CURRENT FIELD NUMBER: PCT/AU00/00988 15 <150> PRIOR APPLICATION NUMBER: PCT/AU00/00988 | |
| 15 <150> PRIOR ETITING DATE: 2000-08-18 | |
| 16 <151> PRIOR FILING DATE: 2000-08-18 | • |
| 16 <151> PRIOR TIPLICATION NUMBER: AU PQ2337 19 <150> PRIOR APPLICATION NUMBER: 1999-08-19 | |
| 20 <151> PRIOR FILING DATE: 1999-08-19 | |
| 23 <160> NUMBER OF SEQ ID NOS: 12 | |
| 26 <170> SOFTWARE: PatentIn version 3.1 | 10 / 11 |
| 29 <210> SEQ ID NO: 1 | r summery sheet, items 10 +11 |
| 30 <211> LENGTH: 27 | n summary of |
| 31 <212> TYPE: DNA Involid reliporte, | |
| 32 (213) ORGINIZES | |
| 35 <400> SEOUENCE: 1 | 27 |
| 36 cgcggatcca gtgctggcat ttgaaga | • |
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| 75 <211> LENGTH: 31 | |
| 76 <212> TYPE: DNA | |
| 77 <213> ORGANISM: Primer | |
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| 80 <4009 SECONICE | , I |
| 81 CGCGGGCCCA GCCCGGCCGGCCGGCGCGGCGGCGGCGGGCG | |
| 84 <210> SEQ ID NO: 6 | |
| 85 <211> LENGTH: 33 | |



PATENT APPLICATION: US/10/049,953

DATE: 03/05/2002 TIME: 14:11:51

Input Set : A:\38247.txt

Output Set: N:\CRF3\03052002\J049953.raw

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/049,953

DATE: 03/05/2002 TIME: 14:11:52

Input Set : A:\38247.txt

Output Set: N:\CRF3\03052002\J049953.raw

L:15 M:270 C: Current Application Number differs, Replaced Current Application No

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date